SEQUENCE LISTING

<110> THE GENERAL HOSPITAL CORPORATION
SHELLEY, CARL SIMON
FAROKHZAD, OMID C.

<120> METHODS FOR DIAGNOSING AND TREATING TUMORS AND SUPPRESSING CD PROMOTERS

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<140> not yet assigned
<141> 2003-09-23

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<151> 2002-09-23

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<170> PatentIn version 3.2

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            20                  25                  30

Leu Val Ser Thr Ser Glu Pro Leu Ser Ser Lys Met Tyr Thr Thr Ser
        35                  40                  45

Ile Thr Ser Asp Pro Lys Ala Asp Ser Thr Gly Asp Gln Thr Ser Ala
    50                  55                  60

Leu Pro Pro Ser Thr Ser Ile Asn Glu Gly Ser Pro Leu Trp Thr Ser
65                  70                  75                  80

Ile Gly Ala Ser Thr Gly Ser Pro Leu Pro Glu Pro Thr Thr Tyr Gln
                85                  90                  95

Glu Val Ser Ile Lys Met Ser Ser Val Pro Gln Glu Thr Pro His Ala
            100                 105                 110

Thr Ser His Pro Ala Val Pro Ile Thr Ala Asn Ser Leu Gly Ser His
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Thr Val Thr Gly Gly Thr Ile Thr Thr Asn Ser Pro Glu Thr Ser Ser
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Ser Arg Gly Thr Ser Gly Pro Pro Leu Thr Met Ala Thr Val Ser Leu
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Glu Thr Ser Lys Gly Thr Ser Gly Pro Pro Val Thr Met Ala Thr Asp
            180                 185                 190

Ser Leu Glu Thr Ser Thr Gly Thr Thr Gly Pro Pro Val Thr Met Thr
        195                 200                 205

Thr Gly Ser Leu Glu Pro Ser Ser Gly Ala Ser Gly Pro Gln Val Ser
    210                 215                 220

Ser Val Lys Leu Ser Thr Met Met Ser Pro Thr Thr Ser Thr Asn Ala
225                 230                 235                 240

Ser Thr Val Pro Phe Arg Asn Pro Asp Glu Asn Ser Arg Gly Met Leu
                245                 250                 255

Pro Val Ala Val Leu Val Ala Leu Leu Ala Val Ile Val Leu Val Ala
            260                 265                 270

Leu Leu Leu Leu Trp Arg Arg Arg Gln Lys Arg Arg Thr Gly Ala Leu
        275                 280                 285

Val Leu Ser Arg Gly Gly Lys Arg Asn Gly Val Val Asp Ala Trp Ala
    290                 295                 300

Gly Pro Ala Gln Val Pro Glu Glu Gly Ala Val Thr Val Thr Val Gly
305                 310                 315                 320

Gly Ser Gly Gly Asp Lys Gly Ser Gly Phe Pro Asp Gly Glu Gly Ser
                325                 330                 335

Ser Arg Arg Pro Thr Leu Thr Thr Phe Phe Gly Arg Arg Lys Ser Arg
            340                 345                 350

Gln Gly Ser Leu Ala Met Glu Glu Leu Lys Ser Gly Ser Gly Pro Ser
        355                 360                 365

Leu Lys Gly Glu Glu Glu Pro Leu Val Ala Ser Glu Asp Gly Ala Val
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<213> Homo sapiens

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<212> PRT

<213> Homo sapiens

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Ala Leu Gly Ser Thr Thr Ala Val Gln Thr Pro Thr Ser Gly Glu Pro
            20                  25                  30

Leu Val Ser Thr Ser Glu Pro Leu Ser Ser Lys Met Tyr Thr Thr Ser
        35                  40                  45

Ile Thr Ser Asp Pro Lys Ala Asp Ser Thr Gly Asp Gln Thr Ser Ala
    50                  55                  60

Leu Pro Pro Ser Thr Ser Ile Asn Glu Gly Ser Pro Leu Trp Thr Ser
65                  70                  75                  80

Ile Gly Ala Ser Thr Gly Ser Pro Leu Pro Glu Pro Thr Thr Tyr Gln
                85                  90                  95

Glu Val Ser Ile Lys Met Ser Ser Val Pro Gln Glu Thr Pro His Ala
            100                 105                 110

Thr Ser His Pro Ala Val Pro Ile Thr Ala Asn Ser Leu Gly Ser His
        115                 120                 125

Thr Val Thr Gly Gly Thr Ile Thr Thr Asn Ser Pro Glu Thr Ser Ser
    130                 135                 140

Arg Thr Ser Gly Ala Pro Val Thr Thr Ala Ala Ser Ser Leu Glu Thr
145                 150                 155                 160

Ser Arg Gly Thr Ser Gly Pro Pro Leu Thr Met Ala Thr Val Ser Leu
                165                 170                 175

Glu Thr Ser Lys Gly Thr Ser Gly Pro Pro Val Thr Met Ala Thr Asp
            180                 185                 190

Ser Leu Glu Thr Ser Thr Gly Thr Thr Gly Pro Pro Val Thr Met Thr
        195                 200                 205

Thr Gly Ser Leu Glu Pro Ser Ser Gly Ala Ser Gly Pro Gln Val Ser
    210                 215                 220

Ser Val Lys Leu Ser Thr Met Met Ser Pro Thr Thr Ser Thr Asn Ala
225                 230                 235                 240

Ser Thr Val Pro Phe Arg Asn Pro Asp Glu Asn Ser Arg Gly Met Leu
                245                 250                 255

Pro Val Ala Val Leu Val Ala Leu Leu Ala Val Ile Val Leu Val Ala
            260                 265                 270

Leu Leu Leu Leu Trp Arg Arg Arg Gln Lys Arg Arg Thr Gly Ala Leu
        275                 280                 285

Val Leu Ser Arg Gly Gly Lys Arg Asn Gly Val Val Asp Ala Trp Ala
    290                 295                 300
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Gly Pro Ala Gln Val Pro Glu Glu Gly Ala Val Thr Val Thr Val Gly
305                 310                 315                 320

Gly Ser Gly Gly Asp Lys Gly Ser Gly Phe Pro Asp Gly Glu Gly Ser
                325                 330                 335

Ser Arg Arg Pro Thr Leu Thr Thr Phe Phe Gly Arg Arg Lys Ser Arg
            340                 345                 350

Gln Gly Ser Leu Ala Met Glu Glu Leu Lys Ser Gly Ser Gly Pro Ser
        355                 360                 365

Leu Lys Gly Glu Glu Glu Pro Leu Val Ala Ser Glu Asp Gly Ala Val
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Asp Ala Pro Ala Pro Asp Glu Pro Glu Gly Gly Asp Gly Ala Ala Pro
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<212> PRT
<213> Homo sapiens leukosialin

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            20                  25                  30

Leu Val Ser Thr Ser Glu Pro Leu Ser Ser Lys Met Tyr Thr Thr Ser
        35                  40                  45

Ile Thr Ser Asp Pro Lys Ala Asp Ser Thr Gly Asp Gln Thr Ser Ala
    50                  55                  60

Leu Pro Pro Ser Thr Ser Ile Asn Glu Gly Ser Pro Leu Trp Thr Ser
65                  70                  75                  80

Ile Gly Ala Ser Thr Gly Ser Pro Leu Pro Glu Pro Thr Thr Tyr Gln
                85                  90                  95

Glu Val Ser Ile Lys Met Ser Ser Val Pro Gln Glu Thr Pro His Ala
            100                 105                 110

Thr Ser His Pro Ala Val Pro Ile Thr Ala Asn Ser Leu Gly Ser His
        115                 120                 125

Thr Val Thr Gly Gly Thr Ile Thr Thr Asn Ser Pro Glu Thr Ser Ser
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Arg Thr Ser Gly Ala Pro Val Thr Thr Ala Ala Ser Ser Leu Glu Thr
145                 150                 155                 160
Ser Arg Gly Thr Ser Gly Pro Pro Leu Thr Met Ala Thr Val Ser Leu
                165                 170                 175

Glu Thr Ser Lys Gly Thr Ser Gly Pro Pro Val Thr Met Ala Thr Asp
            180                 185                 190

Ser Leu Glu Thr Ser Thr Gly Thr Thr Gly Pro Pro Val Thr Met Thr
        195                 200                 205

Thr Gly Ser Leu Glu Pro Ser Ser Gly Ala Ser Gly Pro Gln Val Ser
    210                 215                 220

Ser Val Lys Leu Ser Thr Met Met Ser Pro Thr Thr Ser Thr Asn Ala
225                 230                 235                 240

Ser Thr Val Pro Phe Arg Asn Pro Asp Glu Asn Ser Arg Gly Met Leu
                245                 250                 255

Pro Val Ala Val Leu Val Ala Leu Leu Ala Val Ile Val Leu Val Ala
            260                 265                 270

Leu Leu Leu Leu Trp Arg Arg Arg Gln Lys Arg Arg Thr Gly Ala Leu
        275                 280                 285

Val Leu Ser Arg Gly Gly Lys Arg Asn Gly Val Val Asp Ala Trp Ala
    290                 295                 300

Gly Pro Ala Gln Val Pro Glu Glu Gly Ala Val Thr Val Thr Val Gly
305                 310                 315                 320

Gly Ser Gly Gly Asp Lys Gly Ser Gly Phe Pro Asp Gly Glu Gly Ser
                325                 330                 335

Ser Arg Arg Pro Thr Leu Thr Thr Phe Phe Gly Arg Arg Lys Ser Arg
            340                 345                 350

Gln Gly Ser Leu Ala Met Glu Glu Leu Lys Ser Gly Ser Gly Pro Ser
        355                 360                 365

Leu Lys Gly Glu Glu Glu Pro Leu Val Ala Ser Glu Asp Gly Ala Val
    370                 375                 380

Asp Ala Pro Ala Pro Asp Glu Pro Glu Gly Gly Asp Gly Ala Ala Pro
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<212> PRT
<213> Homo sapiens leukosialin

<400> 8

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Leu Val Ser Thr Ser Glu Pro Leu Ser Ser Lys Met Tyr Thr Thr Ser
        35                  40                  45

Ile Thr Ser Asp Pro Lys Ala Asp Ser Thr Gly Asp Gln Thr Ser Ala
    50                  55                  60

Leu Pro Pro Ser Thr Ser Ile Asn Glu Gly Ser Pro Leu Trp Thr Ser
65                  70                  75                  80

Ile Gly Ala Ser Thr Gly Ser Pro Leu Pro Glu Pro Thr Thr Tyr Gln
                85                  90                  95

Glu Val Ser Ile Lys Met Ser Ser Val Pro Gln Glu Thr Pro His Ala
            100                 105                 110

Thr Ser His Pro Ala Val Pro Ile Thr Ala Asn Ser Leu Gly Ser His
        115                 120                 125

Thr Val Thr Gly Gly Thr Ile Thr Thr Asn Ser Pro Glu Thr Ser Ser
    130                 135                 140

Arg Thr Ser Gly Ala Pro Val Thr Thr Ala Ala Ser Ser Leu Glu Thr
145                 150                 155                 160

Ser Arg Gly Thr Ser Gly Pro Pro Leu Thr Met Ala Thr Val Ser Leu
                165                 170                 175

Glu Thr Ser Lys Gly Thr Ser Gly Pro Pro Val Thr Met Ala Thr Asp
            180                 185                 190

Ser Leu Glu Thr Ser Thr Gly Thr Thr Gly Pro Pro Val Thr Met Thr
        195                 200                 205

Thr Gly Ser Leu Glu Pro Ser Ser Gly Ala Ser Gly Pro Gln Val Ser
    210                 215                 220

Ser Val Lys Leu Ser Thr Met Met Ser Pro Thr Thr Ser Thr Asn Ala
225                 230                 235                 240
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Ser Thr Val Pro Phe Arg Asn Pro Asp Glu Asn Ser Arg Gly Met Leu
                245                 250                 255

Pro Val Ala Val Leu Val Ala Leu Leu Ala Val Ile Val Leu Val Ala
            260                 265                 270

Leu Leu Leu Leu Trp Arg Arg Arg Gln Lys Arg Arg Thr Gly Ala Leu
        275                 280                 285

Val Leu Ser Arg Gly Gly Lys Arg Asn Gly Val Val Asp Ala Trp Ala
    290                 295                 300

Gly Pro Ala Gln Val Pro Glu Glu Gly Ala Val Thr Val Thr Val Gly
305                 310                 315                 320

Gly Ser Gly Gly Asp Lys Gly Ser Gly Phe Pro Asp Gly Glu Gly Ser
                325                 330                 335

Ser Arg Arg Pro Thr Leu Thr Thr Phe Phe Gly Arg Arg Lys Ser Arg
            340                 345                 350

Gln Gly Ser Leu Ala Met Glu Glu Leu Lys Ser Gly Ser Gly Pro Ser
        355                 360                 365

Leu Lys Gly Glu Glu Glu Pro Leu Val Ala Ser Glu Asp Gly Ala Val
    370                 375                 380

Asp Ala Pro Ala Pro Asp Glu Pro Glu Gly Gly Asp Gly Ala Ala Pro
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<213> Homo sapiens leukosialin (CD43)

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<212> PRT
<213> Homo sapiens leukosialin (CD43)

<400> 10

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210 215 220

Ser Val Lys Leu Ser Thr Met Met Ser Pro Thr Thr Ser Thr Asn Ala
225 230 235 240

Ser Thr Val Pro Phe Arg Asn Pro Asp Glu Asn Ser Arg Gly Met Leu
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Gly Pro Ala Gln Val Pro Glu Glu Gly Ala Val Thr Val Thr Val Gly
305 310 315 320

Gly Ser Gly Gly Asp Lys Gly Ser Gly Phe Pro Asp Gly Glu Gly Ser
325 330 335

Ser Arg Arg Pro Thr Leu Thr Thr Phe Phe Gly Arg Arg Lys Ser Arg
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Gln Gly Ser Leu Ala Met Glu Glu Leu Lys Ser Gly Ser Gly Pro Ser
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<211> 1879
<212> DNA
<213> Homo sapiens sialophorin (CD43)

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Ile Thr Ser Asp Pro Lys Ala Asp Ser Thr Gly Asp Gln Thr Ser Ala
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Leu Pro Pro Ser Thr Ser Ile Asn Glu Gly Ser Pro Leu Trp Thr Ser
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Ile Gly Ala Ser Thr Gly Ser Pro Leu Pro Glu Pro Thr Thr Tyr Gln
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Glu Val Ser Ile Lys Met Ser Ser Val Pro Gln Glu Thr Pro His Ala
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Thr Ser His Pro Ala Val Pro Ile Thr Ala Asn Ser Leu Gly Ser His
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Arg Thr Ser Gly Ala Pro Val Thr Thr Ala Ala Ser Ser Leu Glu Thr
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Glu Thr Ser Lys Gly Thr Ser Gly Pro Pro Val Thr Met Ala Thr Asp
180 185 190

Ser Leu Glu Thr Ser Thr Gly Thr Thr Gly Pro Pro Val Thr Met Thr
195 200 205

Thr Gly Ser Leu Glu Pro Ser Ser Gly Ala Ser Gly Pro Gln Val Ser
210 215 220

Ser Val Lys Leu Ser Thr Met Met Ser Pro Thr Thr Ser Thr Asn Ala
225 230 235 240

Ser Thr Val Pro Phe Arg Asn Pro Asp Glu Asn Ser Arg Gly Met Leu
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Pro Val Ala Val Leu Val Ala Leu Leu Ala Val Ile Val Leu Val Ala
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Leu Leu Leu Leu Trp Arg Arg Arg Gln Lys Arg Arg Thr Gly Ala Leu
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Val Leu Ser Arg Gly Gly Lys Arg Asn Gly Val Val Asp Ala Trp Ala
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Gly Ser Gly Gly Asp Lys Gly Ser Gly Phe Pro Asp Gly Glu Gly Ser
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Ser Arg Arg Pro Thr Leu Thr Thr Phe Phe Gly Arg Arg Lys Ser Arg

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gccagaggcc tcacggtttc tcctccgagt ttctggctgg gtgtagttct cagaaacccc 3780
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gtgttaaaag agaggcacag gccgggtgcg gtggctcacg cctgtaatct cagcactttg 5340

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tcccagctac ttaggaggct gaggccagag aattgcttga accctggagt cagaggttgc 5520

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ctgtgcctgg catgttgcag atcctccatg aatatgcatt tgaatgaatg aatgaatgaa 6240

tgaatgaatg gagatgacgc ctcagagatt ctttcttttg agatgaggtc tcattctgtc 6300

acccagacta gagggcagtg gtgcaatcac agctcaccac agcctcaacc tcctgggcct 6360

cccaagtagc tgcgatcaca ggtgtgcacc aacatgccca gctaattttt tttttaattt 6420

ttaatttgta cagacagggt cttgctgtgt tgcccaggct ggtctcgaac tcctgggctc 6480

aagtggtcct cccacctaag ctt 6503

<210> 14
<211> 400
<212> PRT
<213> Homo sapiens sialophorin (CD43)

<400> 14
Met Ala Thr Leu Leu Leu Leu Leu Gly Val Leu Val Val Ser Pro Asp
1 5 10 15

Ala Leu Gly Ser Thr Thr Ala Val Gln Thr Pro Thr Ser Gly Glu Pro
20 25 30

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Leu Val Ser Thr Ser Glu Pro Leu Ser Ser Lys Met Tyr Thr Thr Ser
        35                  40                  45

Ile Thr Ser Asp Pro Lys Ala Asp Ser Thr Gly Asp Gln Thr Ser Ala
    50                  55                  60
Leu Pro Pro Ser Thr Ser Ile Asn Glu Gly Ser Pro Leu Trp Thr Ser
65                  70                  75                  80

Ile Gly Ala Ser Thr Gly Ser Pro Leu Pro Glu Pro Thr Thr Tyr Gln
                85                  90                  95

Glu Val Ser Ile Lys Met Ser Ser Val Pro Gln Glu Thr Pro His Ala
            100                 105                 110

Thr Ser His Pro Ala Val Pro Ile Thr Ala Asn Ser Leu Gly Ser His
        115                 120                 125

Thr Val Thr Gly Gly Thr Ile Thr Thr Asn Ser Pro Glu Thr Ser Ser
    130                 135                 140

Arg Thr Ser Gly Ala Pro Val Thr Thr Ala Ala Ser Ser Leu Glu Thr
145                 150                 155                 160

Ser Arg Gly Thr Ser Gly Pro Pro Leu Thr Met Ala Thr Val Ser Leu
                165                 170                 175

Glu Thr Ser Lys Gly Thr Ser Gly Pro Pro Val Thr Met Ala Thr Asp
            180                 185                 190

Ser Leu Glu Thr Ser Thr Gly Thr Thr Gly Pro Pro Val Thr Met Thr
        195                 200                 205

Thr Gly Ser Leu Glu Pro Ser Ser Gly Ala Ser Gly Pro Gln Val Ser
    210                 215                 220

Ser Val Lys Leu Ser Thr Met Met Ser Pro Thr Thr Ser Thr Asn Ala
225                 230                 235                 240

Ser Thr Val Pro Phe Arg Asn Pro Asp Glu Asn Ser Arg Gly Met Leu
                245                 250                 255

Pro Val Ala Val Leu Val Ala Leu Leu Ala Val Ile Val Leu Val Ala
            260                 265                 270

Leu Leu Leu Leu Trp Arg Arg Arg Gln Lys Arg Arg Thr Gly Ala Leu
        275                 280                 285

Val Leu Ser Arg Gly Gly Lys Arg Asn Gly Val Val Asp Ala Trp Ala
    290                 295                 300

Gly Pro Ala Gln Val Pro Glu Glu Gly Ala Val Thr Val Thr Val Gly
305                 310                 315                 320

Gly Ser Gly Gly Asp Lys Gly Ser Gly Phe Pro Asp Gly Glu Gly Ser
                325                 330                 335

Ser Arg Arg Pro Thr Leu Thr Thr Phe Phe Gly Arg Arg Lys Ser Arg
            340                 345                 350

Gln Gly Ser Leu Ala Met Glu Glu Leu Lys Ser Gly Ser Gly Pro Ser
        355                 360                 365
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Leu Lys Gly Glu Glu Glu Pro Leu Val Ala Ser Glu Asp Gly Ala Val
370 375 380

Asp Ala Pro Ala Pro Asp Glu Pro Glu Gly Gly Asp Gly Ala Ala Pro
385 390 395 400

<210> 15
<211> 2745
<212> DNA
<213> Homo sapiens heterogeneous nuclear ribonucleoprotein K

<400> 15
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cttcagttct gctctgcaag gatatataat aactgattgg tgtgcccgtt taataaaaga 120

atatggaaac tgaacagcca gaagaaacct tccctaacac tgaaaccaat ggtgaatttg 180

gtaaacgccc tgcagaagat atggaagagg aacaagcatt taaaagatct agaaacactg 240

atgagatggt tgaattacgc attctgcttc agagcaagaa tgctggggca gtgattggaa 300

aaggaggcaa gaatattaag gctctccgta cagactacaa tgccagtgtt tcagtcccag 360

acagcagtgg ccccgagcgc atattgagta tcagtgctga tattgaaaca attggagaaa 420

ttctgaagaa aatcatccct accttggaag agggcctgca gttgccatca cccactgcaa 480

ccagccagct cccgctcgaa tctgatgctg tggaatgctt aaattaccaa cactataaag 540

gaagtgactt tgactgcgag ttgaggctgt tgattcatca gagtctagca ggaggaatta 600

ttggggtcaa aggtgctaaa atcaaagaac ttcgagagaa cactcaaacc accatcaagc 660

ttttccagga atgctgtcct cattccactg acagagttgt tcttattgga ggaaaacccg 720

atagggttgt agagtgcata aagatcatcc ttgatcttat atctgagtct cccatcaaag 780

gacgtgcaca gccttatgat cccaattttt acgatgaaac ctatgattat ggtggtttta 840

caatgatgtt tgatgaccgt cgcggacgcc cagtgggatt tcccatgcgg ggaagaggtg 900

gttttgacag aatgcctcct ggtcggggtg ggcgtcccat gcctccatct agaagagatt 960

atgatgatat gagccctcgt cgaggaccac ctccccctcc tcccggacga ggcggccggg 1020

gtggtagcag agctcggaat cttcctcttc ctccaccacc accacctaga gggggagacc 1080

tcatggccta tgacagaaga gggagacctg gagaccgtta cgacggcatg gttggtttca 1140

gtgctgatga aacttgggac tctgcaatag atacatggag cccatcagaa tggcagatgg 1200

cttatgaacc acagggtggc tccggatatg attattccta tgcaggggt cgtggctcat 1260

atggtgatct tggtggacct attattacta cacaagtaac tattcccaaa gatttggctg 1320

gatctattat tggcaaaggt ggtcagcgga ttaaacaaat ccgtcatgag tcgggagctt 1380

cgatcaaaat tgatgagcct ttagaaggat ccgaagatcg gatcattacc attacaggaa 1440

cacaggacca gatacagaat gcacagtatt tgctgcagaa cagtgtgaag cagtatgcag 1500

atgttgaagg attctaatgc aagatatttt ttctttttta tagtgtgaag cagtattctg 1560

gaaagttttt ctaagactag tgaagaactg aaggagtcct gcatcttttt ttttttatct 1620

gcttctgttt aaaaagccaa cattcctctg cttcataggt gttctgcatt tgaggtgtag 1680

tgaaatcttt gctgttcacc agatgtaatg ttttagttcc ttacaaacag ggttgggggg 1740

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tgctttgtgg tcactgtaac atttgggggg tgggacaggg aggaaaagta acaatagtcc 1920

acatgtccct ggcatctgtt cagagcagtg tgcagaatgt aatgctcttt tgtaagaaac 1980

gttttatgat ttttaaaata aatttagtga acctatttt ggtggtcatt tttttttaa 2040

gacagtcatt ttaaaatggt ggctgaattt cccaacccac ccccaaacta aacactaagt 2100

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aattccatca tacatattca tgcttataat aagctgggga ttttttgttt gtttttgcaa 2280

atgcttgccc ctacttttca acaattttct atgttagttg tgaagaacta aggtggggag 2340

cagtactaca agttgagtaa tggtatgagt atataccaga attctgattg gcagcaagtt 2400

tattaatcag aataacactt ggttatggaa gtgactaatg ctgaaaaaat tgattatttt 2460

tattagataa tttctcacct atagacttaa actgtcaatt tgctctagtg tcttattagt 2520

taaactttgt aaaatatata tatacttgtt tttccattgt atgcaaattg aaagaaaaag 2580

atgtaccatt tctctgttgt atgttggatt atgtaggaat gtttgtgtac aattcaaaaa 2640

aaaaaaagat gaaaaaagtt cctgtggatg ttttgtgtag tatcttggca tttgtattga 2700

tagttaaaat tcacttccaa ataaataaaa cacccatgat gctag 2745

<210> 16
<211> 463
<212> PRT
<213> Homo sapiens heterogeneous nuclear ribonucleoprotein complex K

<400> 16
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Gly Glu Phe Gly Lys Arg Pro Ala Glu Asp Met Glu Glu Glu Gln Ala
20 25 30

Phe Lys Arg Ser Arg Asn Thr Asp Glu Met Val Glu Leu Arg Ile Leu
35 40 45

Leu Gln Ser Lys Asn Ala Gly Ala Val Ile Gly Lys Gly Gly Lys Asn
50 55 60

Ile Lys Ala Leu Arg Thr Asp Tyr Asn Ala Ser Val Ser Val Pro Asp
65 70 75 80

Ser Ser Gly Pro Glu Arg Ile Leu Ser Ile Ser Ala Asp Ile Glu Thr
85 90 95

Ile Gly Glu Ile Leu Lys Lys Ile Ile Pro Thr Leu Glu Glu Gly Leu
100 105 110

Gln Leu Pro Ser Pro Thr Ala Thr Ser Gln Leu Pro Leu Glu Ser Asp
115 120 125

Ala Val Glu Cys Leu Asn Tyr Gln His Tyr Lys Gly Ser Asp Phe Asp
130 135 140

Cys Glu Leu Arg Leu Leu Ile His Gln Ser Leu Ala Gly Gly Ile Ile
145 150 155 160

Gly Val Lys Gly Ala Lys Ile Lys Glu Leu Arg Glu Asn Thr Gln Thr
165 170 175

Thr Ile Lys Leu Phe Gln Glu Cys Cys Pro His Ser Thr Asp Arg Val
180 185 190

Val Leu Ile Gly Gly Lys Pro Asp Arg Val Val Glu Cys Ile Lys Ile
195 200 205

Ile Leu Asp Leu Ile Ser Glu Ser Pro Ile Lys Gly Arg Ala Gln Pro
210 215 220

Tyr Asp Pro Asn Phe Tyr Asp Glu Thr Tyr Asp Tyr Gly Gly Phe Thr
225 230 235 240

Met Met Phe Asp Asp Arg Arg Gly Arg Pro Val Gly Phe Pro Met Arg
245 250 255

Gly Arg Gly Gly Phe Asp Arg Met Pro Pro Gly Arg Gly Gly Arg Pro
260 265 270

Met Pro Pro Ser Arg Arg Asp Tyr Asp Asp Met Ser Pro Arg Arg Gly
275 280 285

Pro Pro Pro Pro Pro Pro Gly Arg Gly Gly Arg Gly Gly Ser Arg Ala
290 295 300

Arg Asn Leu Pro Leu Pro Pro Pro Pro Pro Pro Arg Gly Gly Asp Leu
305 310 315 320

Met Ala Tyr Asp Arg Arg Gly Arg Pro Gly Asp Arg Tyr Asp Gly Met
325 330 335

Val Gly Phe Ser Ala Asp Glu Thr Trp Asp Ser Ala Ile Asp Thr Trp
340 345 350

Ser Pro Ser Glu Trp Gln Met Ala Tyr Glu Pro Gln Gly Gly Ser Gly
355 360 365

Tyr Asp Tyr Ser Tyr Ala Gly Gly Arg Gly Ser Tyr Gly Asp Leu Gly
370 375 380

Gly Pro Ile Ile Thr Thr Gln Val Thr Ile Pro Lys Asp Leu Ala Gly
385 390 395 400

Ser Ile Ile Gly Lys Gly Gly Gln Arg Ile Lys Gln Ile Arg His Glu
405 410 415

Ser Gly Ala Ser Ile Lys Ile Asp Glu Pro Leu Glu Gly Ser Glu Asp
420 425 430
Arg Ile Ile Thr Ile Thr Gly Thr Gln Asp Gln Ile Gln Asn Ala Gln
435 440 445

Tyr Leu Leu Gln Asn Ser Val Lys Gln Tyr Ser Gly Lys Phe Phe
450 455 460

<210> 17
<211> 1144
<212> DNA
<213> Homo sapiens Pur (pur-alpha)

<400> 17
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tggcggaccg agacagcggc agcgagcagg gtggtgcggc gctgggttcg ggcggctccc 120

tggggcaccc cggctcgggc tcaggctccg gcgggggcgg tggtggcggc gggggcggcg 180

gcggcagtgg cggcggcggc ggcggggccc caggggggct gcagcacgag acgcaggagc 240

tggcctccaa gcgggtggac atccagaaca agcgcttcta cctggacgtg aagcagaacg 300

ccaagggccg cttcctgaag atcgccgagg tgggcgcggg cggcaacaag agccgcctta 360

ctctctccat gtcagtggcc gtggagttcc gcgactacct gggcgacttc atcgagcact 420

acgcgcagct gggccccagc cagccgccgg acctggccca ggcgcaggac gagccgcgcc 480

gggcgctcaa aagcgagttc ctggtgcgcg agaaccgcaa gtactacatg gatctcaagg 540

agaaccagcg cggccgcttc ctgcgcatcc gccagacggt caaccggggg cctggcctgg 600

gctccacgca gggccagacc attgcgctgc ccgcgcaggg gctcatcgag ttccgtgacg 660

ctctggccaa gctcatcgac gactacggag tggaggagga gccggccgag ctgcccgagg 720

gcacctcctt gactgtggac aacaagcgct tcttcttcga tgtgggctcc aacaagtacg 780

gcgtgtttat gcgagtgagc gaggtgaagc ccacctatcg caactccatc accgtgccct 840

acaaggtgtg ggccaagttc ggacacacct tctgcaagta ctcggaggag atgaagaaga 900

ttcaagagaa gcagagggag aagcgggctg cctgtgagca gcttcaccag cagcaacagc 960

agcagcagga ggagaccgcc gctgccactc tgctactgca gggtgaggaa gaaggggaag 1020

aagattgatc aaacagaatg aaacccccac acacacacac atgcatacac acacacacac 1080

agccacacac acagaaaata tactgtaaag aaagagagaa aataaaaagt taaaaagtta 1140

aaaa 1144

<210> 18
<211> 322
<212> PRT

<213> Homo sapiens purine-rich element binding protein A (PURA)

<400> 18

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Met Ala Asp Arg Asp Ser Gly Ser Glu Gln Gly Gly Ala Ala Leu Gly
1               5                  10                  15

Ser Gly Gly Ser Leu Gly His Pro Gly Ser Gly Ser Gly Ser Gly Gly
            20                  25                  30

Gly Gly Gly Gly Gly Gly Gly Gly Gly Gly Ser Gly Gly Gly Gly Gly
        35                  40                  45

Gly Ala Pro Gly Gly Leu Gln His Glu Thr Gln Glu Leu Ala Ser Lys
    50                  55                  60

Arg Val Asp Ile Gln Asn Lys Arg Phe Tyr Leu Asp Val Lys Gln Asn
65                  70                  75                  80

Ala Lys Gly Arg Phe Leu Lys Ile Ala Glu Val Gly Ala Gly Gly Asn
                85                  90                  95

Lys Ser Arg Leu Thr Leu Ser Met Ser Val Ala Val Glu Phe Arg Asp
            100                 105                 110

Tyr Leu Gly Asp Phe Ile Glu His Tyr Ala Gln Leu Gly Pro Ser Gln
        115                 120                 125

Pro Pro Asp Leu Ala Gln Ala Gln Asp Glu Pro Arg Arg Ala Leu Lys
    130                 135                 140

Ser Glu Phe Leu Val Arg Glu Asn Arg Lys Tyr Tyr Met Asp Leu Lys
145                 150                 155                 160

Glu Asn Gln Arg Gly Arg Phe Leu Arg Ile Arg Gln Thr Val Asn Arg
                165                 170                 175

Gly Pro Gly Leu Gly Ser Thr Gln Gly Gln Thr Ile Ala Leu Pro Ala
            180                 185                 190

Gln Gly Leu Ile Glu Phe Arg Asp Ala Leu Ala Lys Leu Ile Asp Asp
        195                 200                 205

Tyr Gly Val Glu Glu Glu Pro Ala Glu Leu Pro Glu Gly Thr Ser Leu
    210                 215                 220

Thr Val Asp Asn Lys Arg Phe Phe Phe Asp Val Gly Ser Asn Lys Tyr
225                 230                 235                 240

Gly Val Phe Met Arg Val Ser Glu Val Lys Pro Thr Tyr Arg Asn Ser
                245                 250                 255

Ile Thr Val Pro Tyr Lys Val Trp Ala Lys Phe Gly His Thr Phe Cys
            260                 265                 270

Lys Tyr Ser Glu Glu Met Lys Lys Ile Gln Glu Lys Gln Arg Glu Lys
        275                 280                 285

Arg Ala Ala Cys Glu Gln Leu His Gln Gln Gln Gln Gln Gln Gln Glu
    290                 295                 300

Glu Thr Ala Ala Ala Thr Leu Leu Leu Gln Gly Glu Glu Glu Gly Glu
305                 310                 315                 320
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Glu Asp

<210> 19
<211> 22
<212> DNA
<213> Synthetic oligonucleotide (CD43 PyRo SS)

<400> 19
gggcccactt cctttccccct tg 22

<210> 20
<211> 16
<212> DNA
<213> Synthetic oligonucleotide (CD43 PyRo SSUB)

<220>
<221> misc_feature
<222> (9)..(10)
<223> bromouracil

<220>
<221> misc_feature
<222> (13)..(15)
<223> bromouracil

<220>
<221> misc_feature
<222> (20)..(21)
<223> bromouracil

<220>
<221> misc_feature
<222> (23)..(23)
<223> biotin

<400> 20
gggcccaccc ccccgb 16

<210> 21
<211> 22
<212> DNA
<213> Synthetic oligonucleotide (CD43 Mut-11)

<400> 21
gggcccactt ccttcatata tg 22

<210> 22
<211> 20
<212> DNA
<213> Synthetic oligonucleotide (NS-SS)

<400> 22
gagttagctc actcattagg 20

<210> 23
<211> 21
<212> DNA
<213> Synthetic oligonucleotide(LUC-2)

<400> 23
atagccttat gcagttgctc t 21

<210> 24
<211> 39
<212> DNA
<213> Synthetic oligonucleotide (GeneRacer RNA Oligo)

<220>
<221> misc_feature
<222> (5)..(5)
<223> bromouracil

<220>
<221> misc_feature
<222> (21)..(21)
<223> bromouracil

<220>
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<222> (26)..(26)
<223> bromouracil

<220>
<221> misc_feature
<222> (31)..(31)
<223> bromouracil

<220>
<221> misc_feature
<222> (39)..(39)
<223> bromouracil

<400> 24
cgacggagca cgaggacacg acaggacgaa ggagagaaa 39

<210> 25
<211> 54
<212> DNA
<213> Synthetic oligonucleotide (GeneRacer Oligo dT Primer)

<400> 25
gctgtcaacg atacgctacg taacggcatg acagtgtttt ttttttttt tttt 54

<210> 26
<211> 23
<212> DNA
<213> Synthetic oligonucleotide (GeneRacer 5' Primer)

<400> 26
cgactggagc acgaggacac tga 23

<210> 27
<211> 27
<212> DNA
<213> Synthetic oligonucleotide (GeneRacer 5′ Nested Primer)

<400> 27
ggacactgac catggactga aggagta 27

<210> 28
<211> 33
<212> DNA
<213> Synthetic oligonucleotide (LUC-4)

<400> 28
cactacggta ggctgcgaaa tgttcatact gtt 33